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QTDNNLIDEFSFQTPTMSTLDLTQNPTVDKVNEHAPTYINTSPNKSIMKKATPK
ASPKKVAFTVTNPEIHHYPDNRVEEEDQSQQKEDSVEPPLIQHQWKDPSQFNYS
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SDEVVESLLPRDLSRDKLETTKEHDAPEHNNENFIDAKSTNTKGQLLVSSDDHL
DSFDRSYNHTEQSILNLLNSASQSQISLNNALEKQRQTQEQQEQTQAAEPEEETSF
DNIKVQEPKSNLEFVKVTIKKEPVSAIEKAPKREFSSRILRIKNEDEIAEPADIHP
KKENEANSHVEDTDALLKKALNDDEESDTTQNSTKMSIRFHIDSDWKLEDSNDG
DREDNDDISRFEKSDILNDVSQTSIDIIGDKYGNSSSEITTCTLAPPERSDNNDKENS
KSLEDPANNESLQQQLEVPHTKEDDSILANSSNIAPPEELTPVVEANDYSSFND
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YKNEQESRVTSKVKIPNAIQFKKFKEVNMSRRV/SPDMDDLNVSQFLPELSE
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QKPKKNSIVTDPEDRYEELQQTASIHNATIDSSIYGRPDSISTDMLPYLSDELKKP
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NKDVQHKPREKQKQKHHHRHHHHHKQKTDIPGVVDEIPDVLQERGKLFFR
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EFILTLKASYEKPRGLTVEVTEKKV/KSRNRLSRLFGSKDIITTTKFVPTEVKDTWA
NKFAPDGSFARCYIDLQQFEDQITGKASQFDLNCFNEWETMSNGNQPMKRGKP
YKIAQLEVKMLYVPRSDPREILPTSIRSAYESINELNNEQNNYFEGYLHQEGGDC
PIFKRFFKLMGTSLLAHSEISHKTRAKINLSKVVDLIYVDKENIDRSNHRNFSDVL
LLDHAFKIKFANGELIDFCAPNKHEMKIWIQNLQEIIYRNRFRRQPWVNMLQQQ
QQQQQQQSSQQ

FIGURE 1

1 cccaaaaaaag ataaaataaa aacaaaacaa aacaaaagta ctaacaaaatt attgaaact
61 ttaatttta ataaaagaatc agtagatcta ttgtaaaag aatgaactc aactccaagt
121 aaatttattac cgatagataa acattctcat ttacaattac agcctcaatc gtcctcgca
181 tcaatattt attccccaaac aaaaccattt aattccccaaac gaacaaaattc caagccgagtt
241 ttagatccaa attcaagctc tgatacctac actagcgaac aagatcaaga gaaagggaaa
301 gaagagaaaaa aggacacacgc cttaacaaca tctttgata gaaatttga tcttgataat
361 tcaatcgata tacaacaaac aattcaacat cagcaacaac agccacaaca acaacaacaa
421 ctctcacaaa ccgacaataa ttaattgtat gaattttctt ttcaaacacc gatgactcg
481 acttttagacc taaccaagca aatccaact gtggacaaag tgaatgaaaaa tcatgcacca
541 actttatataa atacccccc caacaaatca ataatgaaaaa aggcaactcc taaagcgta
601 cctaaaaaaag ttgcatttac ttaactaat cccgaaatc atcaattatcc agataataga
661 gtcgaggaag aagatcaaag tcaacaaaaa gaagattcag ttgagccacc cttatataa
721 catcaatgaa aagatcctc tcaattcaat tattctgatg aagatacaaa tgctcagtt
781 ccaccaacac caccactca tacgacgaaa cctactttt cgcaatttattt gaacaaaaac
841 aacgaagtca atctggacc agaggcattt acagatatga aattaaagcg cgaaaaattt
901 agcaatttat cattagatga aaaagtcaat ttatatctt gtcccactaa tataatacaat
961 agtaagaatg tgcagatattt ggatctgcat ttacaaaact tgcaagacgc ttggaaaaac
1021 aaaactaatg aaaatattca caatttttca ttgcattttaa aagcaccaaa gaatgatatt
1081 gaaaacccat taaactcatt gactaacgca gatattctgt taagatcatc tggatcatca
1141 caatcgatcat tacaatctt gaggaatgac aatctgtct tggatcagt gcctgggtca
1201 cctaagaagg ttaatctgg attgtcttgc aatgacggca taaaggggtt ctctgatgag
1261 gttgtgaat cattacttcc tctgtactt tctcgagaca aatttagagac tacaacaaaa
1321 catgatgcac cagaacacaa caatgagaat ttattgtat ctaatcgac taataccat
1381 aagggacaac tcttagtac atctgtatgat cattttggact cttttgatag atccataac
1441 cacactgaac aatcaattttaa gaatctttt aatagtgcatacataatc aatttcgttta
1501 aatgcattgg aaaaacaaag gcaaacacag gaacaagaac aaacacaacgc ggcagagcc
1561 gaagaagaaaaa cttcgatgt tgataatatc aaagttaaac aagagccaaa gagcaattt
1621 gagttgtca aggttaccat caagaaagaaa ccagttctgg ccacggaaat aaaagctcca
1681 aaaagagaat ttcaagtcg aatattaaga ataaaaaaatg aagatgaaat tgccgaacca
1741 gctgatattc atccaaaaaa agaaaaatgaa gcaaacagtc atgtcaaga tactgtatca
1801 ttgtgtgaaga aagcacttaa tgatgtatgaa gatctgaca cgacccaaaa ctcacgaaa
1861 atgtcaatttgc ttttcatat tgatgtatgat tgaaaattgg aagacagttaa tgatggcgat
1921 agagaagata atgtatgat ttctcgatgtt gggaaatcag atatttgaa cgacgtatca
1981 cagacttctg atattttgg tgacaaatat gggaaactcat caagtggaaat aaccacccaaa
2041 acattagcac ccccaagatc ggacaacaat gacaaggaga attctaaatc ttggaaagat
2101 ccagctataa atgaatcattt gcaacaacaa ttggaggatc cgcatacaaa agaagatgat
2161 agcatttttag ccaactcgatc caatattgtt ccacccgttggaaat gcccgtatgt
2221 gaagcaaaatg attattcatc tttaatgtatc gtgacccaaa ctttgtatgc atactcaacgc
2281 ttgtgtgtatc cttatcttag agagcacgaa actgtatca aaccaattaa ttcatatca
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2401 atcattgtatc gttatcaaca atacaacaaac gaacaagaat ctctgtatc tagtgataaa
2461 gtggaaaatcc caaatggccat acaattcaag aaattcaag aggttttttttcc agaattatct
2521 agagttgtta gtccagacat ggtatgttgc aatgtatctc aatttttacc agaattatct
2581 gaagactctg gatttaaaga ttgtatcc gccaactact ccaataacac caacagacca
2641 agaagttta ctccatgtatc cactaaaaat gtctgtatc atattgtatcaat cgatcataat

FIGURE 2A

FIGURE 2B

Activation of "Subtilisin-like" Proprotein Convertases

Signal peptide	<u>Propeptide</u> Xn-K/R	Inactive Subtilisin D H N S	<u>P-Domain</u> D - H - N <RGD> S Substrate = K/R↓
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The processing or "P-domain" clips the propeptide at the carboxy terminal side of dibasic residues, thereby releasing the propeptide. Exposed D - H - N - S active site residues assume the subtilisin serine protease conformation.

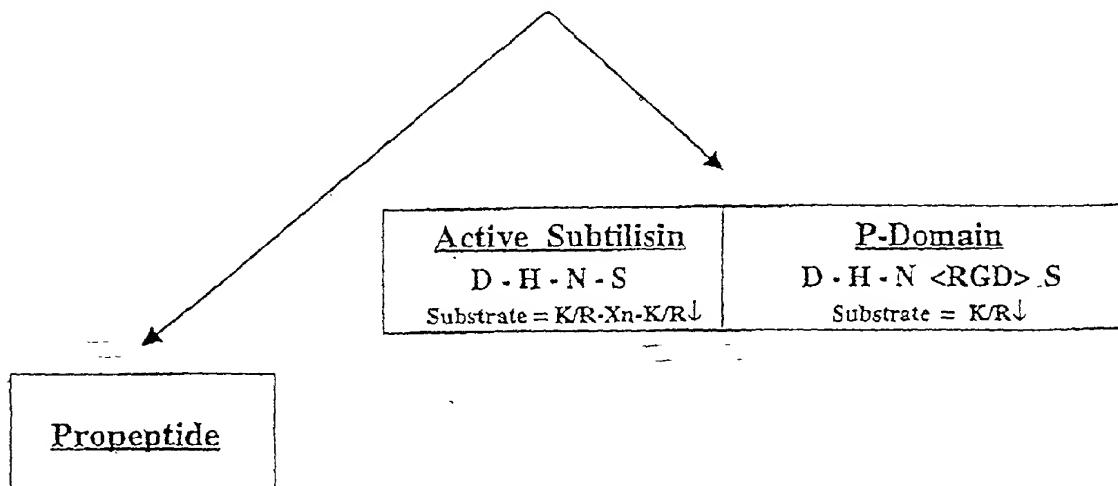
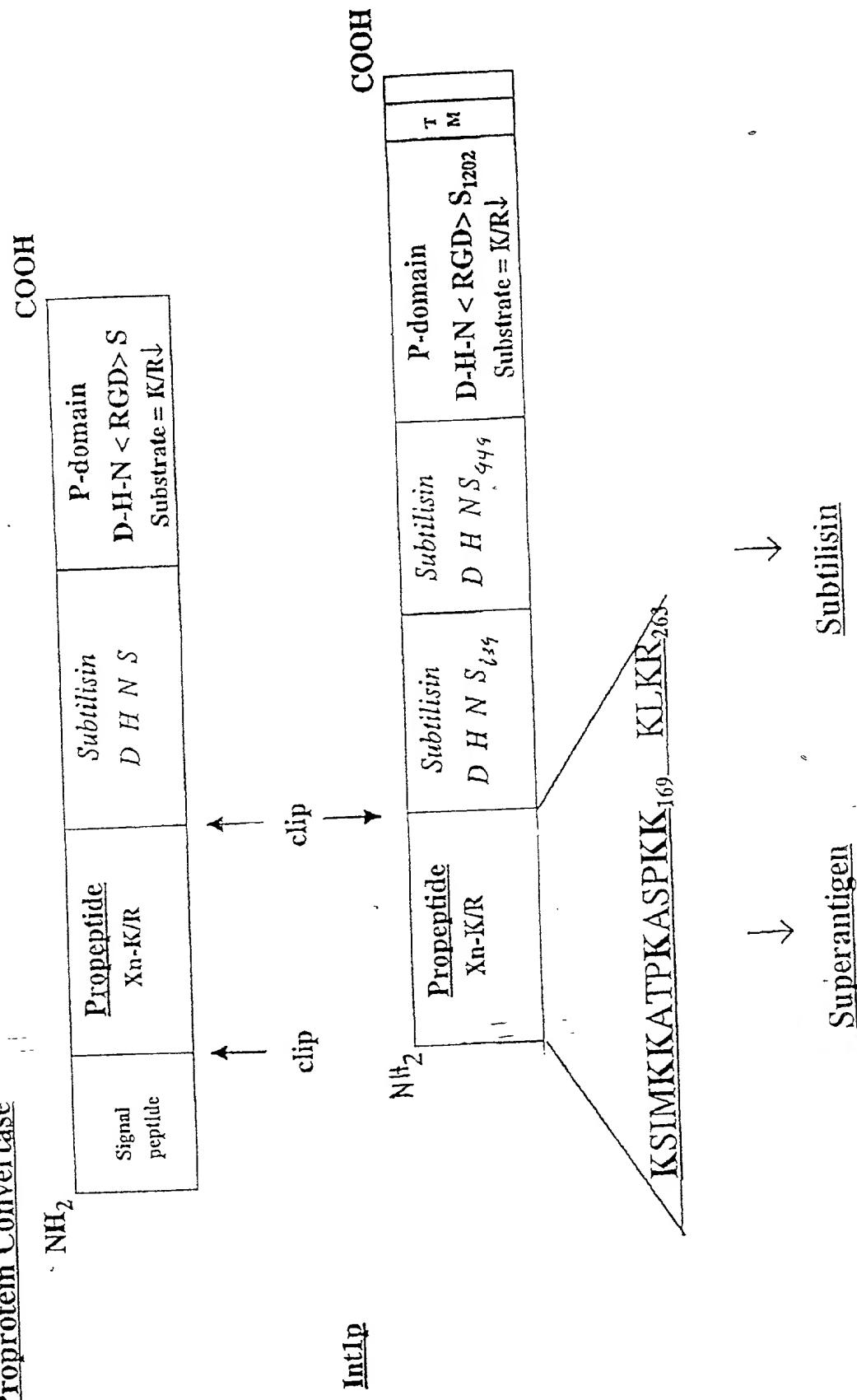


FIG. 3

Amino terminal processing of Int1p

Proprotein Convertase



P Domain Subtilisin Motifs

Kex2 D₁₇₉ H₂₁₃ N₃₁₄ S₃₇₈ = 199aa
 < R₃₁₈GD >

Furin D₃₅₅ H₃₉₅ N₄₇₉ S₅₅₅ = 200aa
 < R₄₉₈GD >

Int1p D₁₀₂₂ H₁₀₆₄ N₁₁₄₆ S₁₂₃₄ = 215aa
 < R₁₁₄₉GD >

CD18 D₂₉₀ H₃₀₉ N₃₅₁ S₄₉₀ = 200aa
 < R₃₉₇GD >

C3 D₁₂₄₅ H₁₂₈₉ N₁₃₂₇ S₁₄₃₀ = 185aa
 < R₁₃₉₃GD >

SpeB D₁₃₅ H₁₅₉ N₂₉₅ S₃₂₄ = 189aa
 < R₃₀₇GD >

Fibrillin D₉₃₀ H₉₇₁ N₁₀₅₂ S₁₁₂₉ = 199aa
 < R₁₀₅₃GD >

EGF D₂₁₉ H₂₈₆ N₃₁₂ S₄₀₃ = 184aa
 < R₃₆₃GD >

Fibronectin D₁₃₆₅ H₁₃₉₆ N₁₄₈₈ S₁₅₆₅ = 200aa
 < R₁₅₆₅GD >

FIG. 5

Comparison of the high affinity heparin-binding site of *Mycobacterium tuberculosis* heparin-binding hemagglutinin adhesin (HBHA) with the proposed heparin-binding site of *Candida albicans* Int1p

HBHA K₁₈₀ AAA KK APA KK AAA KK₁₉₅

Int1p K₁₅₅ SIM KK ATP K ASP KK₁₆₉

FIG. 6

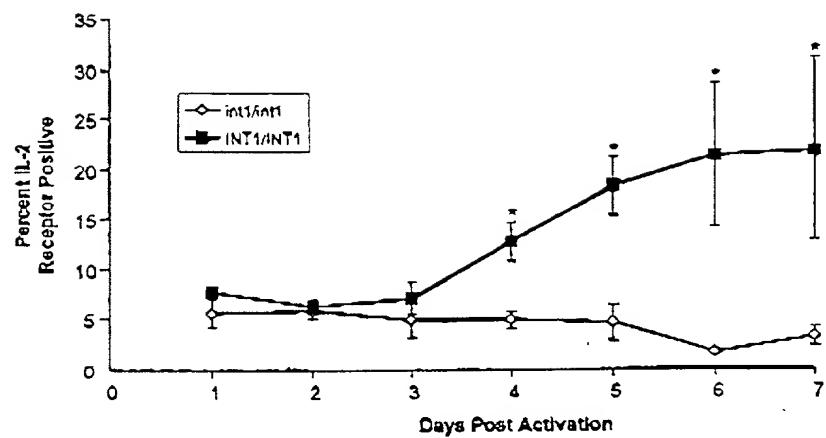


FIG. 7

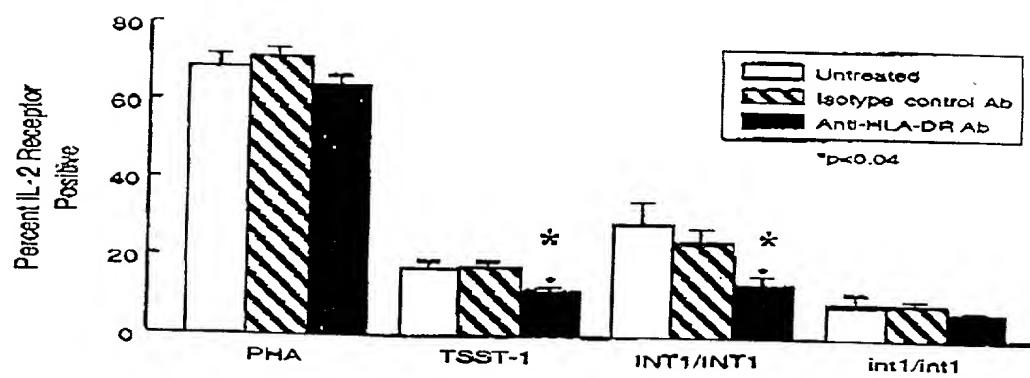


FIG. 8

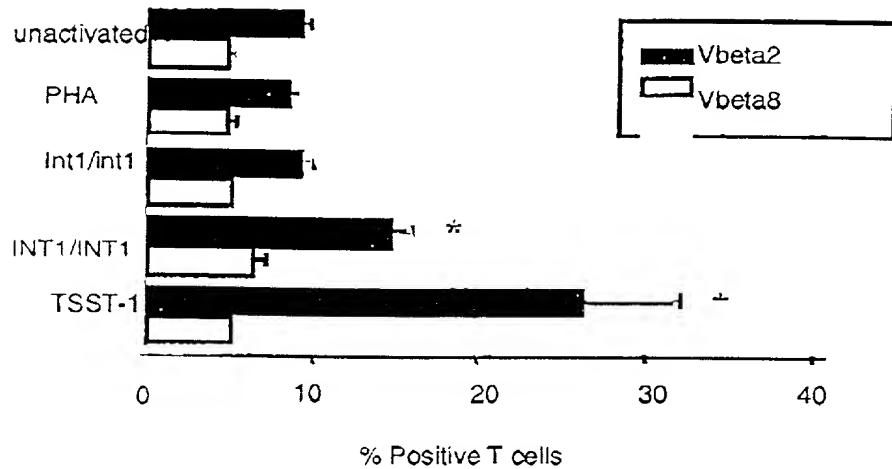


FIG. 9

Si g n al	PRO- PEPTIDE KR	CATALYTIC DOMAIN D(DX)-H-N-S	PROCESSING DOMAIN D-H-N-RGD-S	C-TERMINAL EXTENSION
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▲

FIG. 10

[anti- <i>CB52</i>]		[anti-RGD]		
PRO- PEPTIDE KR	"CATALYTIC DOMAIN 1" D(DX)-H-N-S	"CATALYTIC DOMAIN 2" D(DX)-H-N-S	"PROCESSING DOMAIN" D-H-N-RGD-S	C-TERMINAL EXTENSION
1 263	435	639	738 949 1022	1236 1664

▲ anti-INT600

FIG. 11

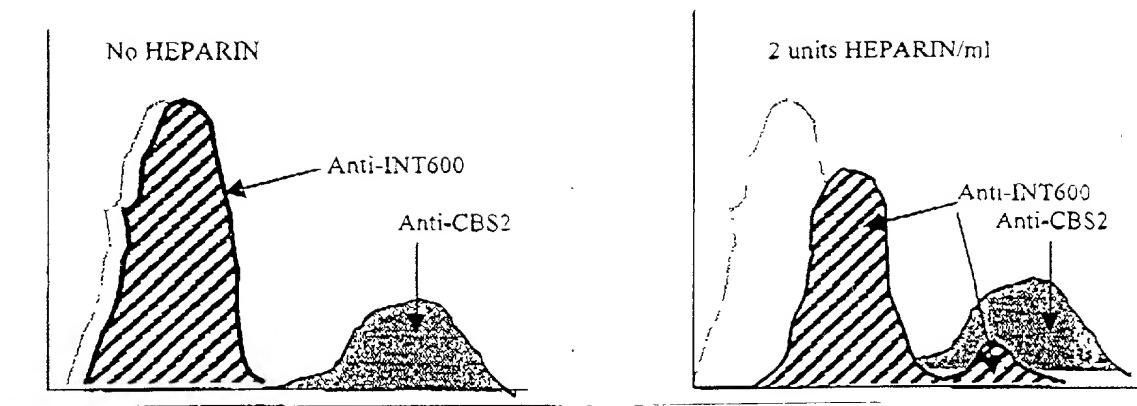


FIG. 12

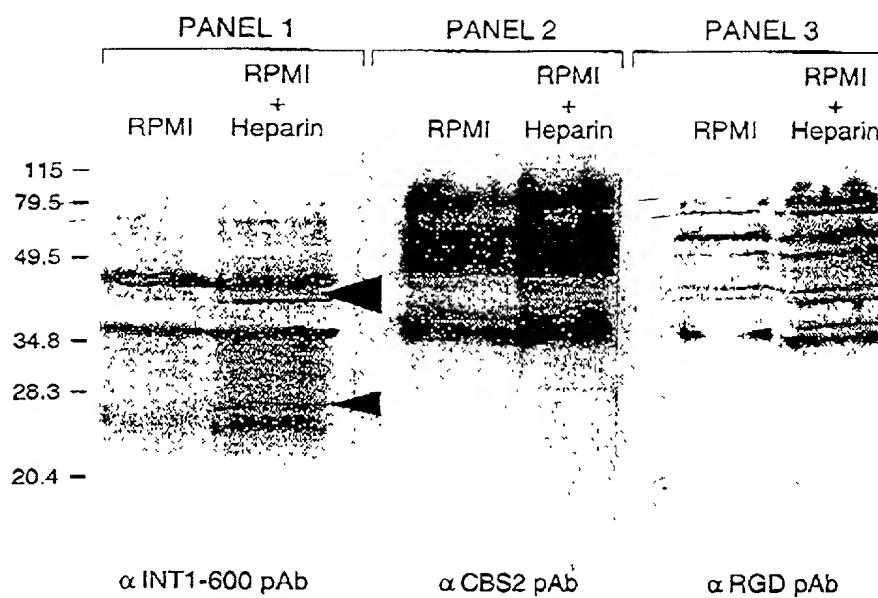


FIG. 13

SILVER STAIN

Anti 6X His WESTERN

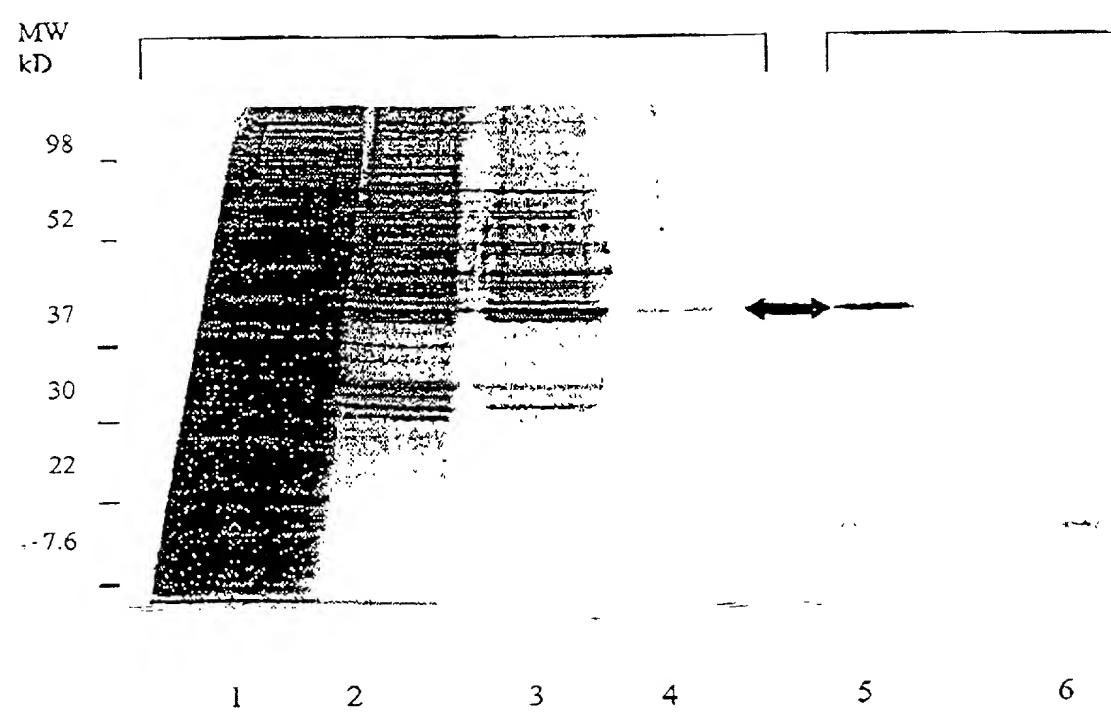


FIG. 14

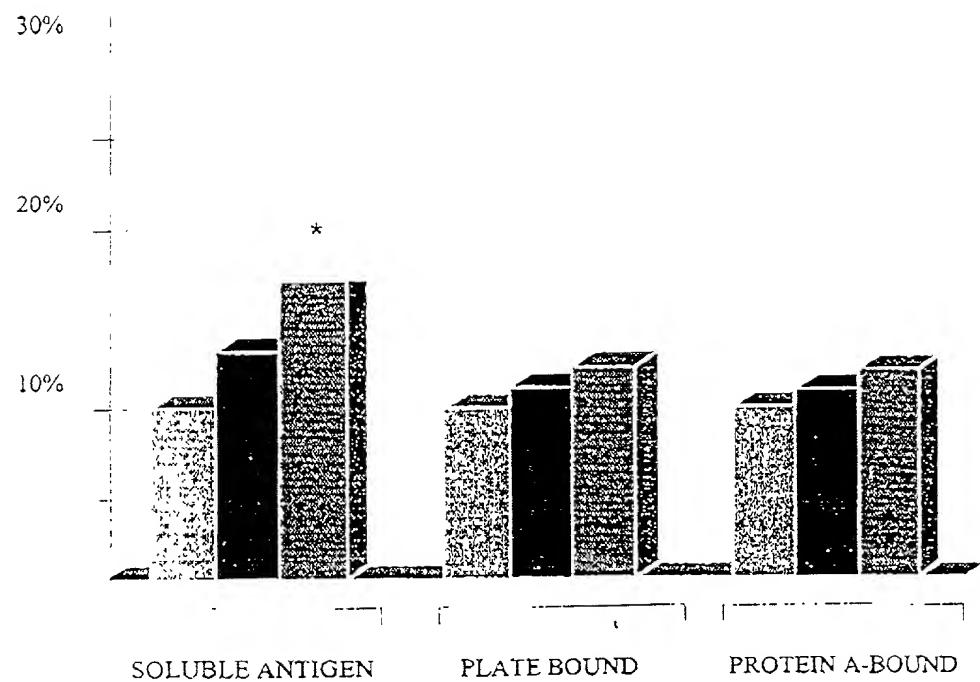


FIG 15

Model for the Participation of Int1p in Candidemia

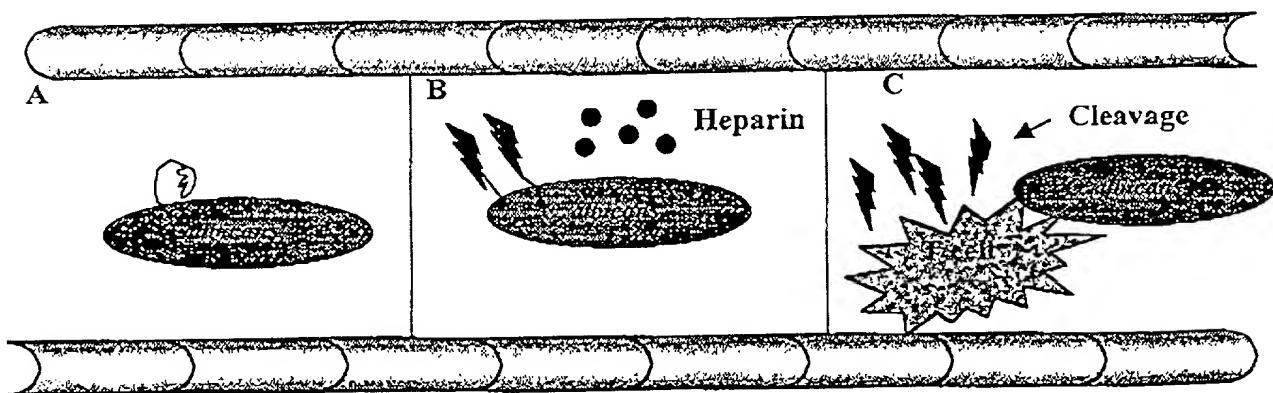


FIG. 16

MHC Class II-Binding Peptides

MAM 15 F V Q N L - - N N V V F T N K E L E₃₁
Intlp 239 F A Q L N K N N E V - - N S E P E₂₅₄

FIG. 17

FIG. 18

Linkage of T Lymphocyte to Antigen-Presenting Cell

